

SEQUENCE LISTING <110> Lu, Shan Shen, Siyuan <120> COMPOSITIONS AND METHODS FOR CYTOMEGALOVIRUS TREATMENT <130> 07917-190001 <140> US 10/789,247 <141> 2004-02-27 <150> US 60/450,818 <151> 2003-02-27 <160> 16 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 2721 <212> DNA <213> Human cytomegalovirus <220> <221> CDS <222> (1)...(2718) <400> 1 atg gaa too agg ato tgg tgc ctg gta gto tgc gtt aac ctg tgt atc 48 Met Glu Ser Arg Ile Trp Cys Leu Val Val Cys Val Asn Leu Cys Ile 10 96 Val Cys Leu Gly Ala Ala Val Ser Ser Ser Ser Thr Ser His Ala Thr 20 25 30 tot tot act cac aat gga ago cat act tot ogt acg acg tot got caa 144 Ser Ser Thr His Asn Gly Ser His Thr Ser Arg Thr Thr Ser Ala Gln 35 acc cgg tca gtc tat tct caa cac gta acg tct tct gaa gcc gtc agt 192 Thr Arg Ser Val Tyr Ser Gln His Val Thr Ser Ser Glu Ala Val Ser 50 55 cat aga gcc aac gag act atc tac aac act acc ctc aag tac gga gat 240 His Arg Ala Asn Glu Thr Ile Tyr Asn Thr Thr Leu Lys Tyr Gly Asp 65 80 gtg gtg gga gtc aac act acc aag tac ccc tat cgc gtg tgt tct atg 288 Val Val Gly Val Asn Thr Thr Lys Tyr Pro Tyr Arg Val Cys Ser Met gcc cag ggt acg gat ctt att cgc ttt gaa cgt aat atc atc tgc acc 336 Ala Gln Gly Thr Asp Leu Ile Arg Phe Glu Arg Asn Ile Ile Cys Thr

		aag Lys 115														384
	-	cgc Arg														432
_	-	ttg Leu	_		_	_	_		_							480
-	_	ggc Gly	_		_	_						_				528
		atc Ile		_		-		-		-			-	-	_	576
		ggc Gly 195	_	-		_	-				_	_		-		624
		atg Met														672
		gtg Val														720
		cgt Arg					-		-	-	_					768
	_	tcc Ser	_							-			_		-	816
	_	tac Tyr 275										_		_	-	864
		gga Gly	-		_	_	_					_				. 912
	-	tcc Ser	_			_			_			-				960
		gct Ala			_	_	_	_	_					_		1008

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	gac Asp															1056
	cgt Arg															1104
	aaa Lys 370															1152
	gac Asp															1200
	cag Gln															1248
	gtg Val															1296
	atc Ile															1344
	agt Ser 450															1392
	aca Thr															1440
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	ctg Leu															1536
	gag Glu															1584
	gcc Ala 530															1632
	ggc Gly															1680
ctg	cgt	gat	atg	aac	gtg	aag	gaa	tcg	сса	gga	cgc	tgc	tac	tca	cga	1728

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				Glu	gac Asp												1824	
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					ttt Phe												2064	
					gta Val												2112	
÷					aag Lys												2160	
					gtt Val 725												2208	
					gaa Glu												2256	
					atc Ile												2304	
					cga Arg												2352	
					ctg Leu												2400	

785	790	795	800
agc acc aaa gac a Ser Thr Lys Asp T 8		_	
gtt tat aat tct g Val Tyr Asn Ser G 820			
tcc acg gcg gct c Ser Thr Ala Ala P 835		Asn Glu Gln Ala	
ctg gcc ctg gcc c Leu Ala Leu Ala A 850			
aca gat tct ttg g Thr Asp Ser Leu A 865			
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Leu Leu Gly Ser Asn Thr Glu Tyr Val Ala Pro Pro Met Trp Glu Ile 170 His His Ile Asn Lys Phe Ala Gln Cys Tyr Ser Ser Tyr Ser Arg Val 185 Ile Gly Gly Thr Val Phe Val Ala Tyr His Arg Asp Ser Tyr Glu Asn 200 205 Lys Thr Met Gln Leu Ile Pro Asp Asp Tyr Ser Asn Thr His Ser Thr 215 Arg Tyr Val Thr Val Lys Asp Gln Trp His Ser Arg Gly Ser Thr Trp 230 235 Leu Tyr Arg Glu Thr Cys Asn Leu Asn Cys Met Leu Thr Ile Thr Thr 245 250 Ala Arg Ser Lys Tyr Pro Tyr His Phe Phe Ala Thr Ser Thr Gly Asp 260 265 Val Val Tyr Ile Ser Pro Phe Tyr Asn Gly Thr Asn Arg Asn Ala Ser 280 Tyr Phe Gly Glu Asn Ala Asp Lys Phe Phe Ile Phe Pro Asn Tyr Thr 295 300 Ile Val Ser Asp Phe Gly Arg Pro Asn Ala Ala Pro Glu Thr His Arg 310 315 Leu Val Ala Phe Leu Glu Arg Ala Asp Ser Val Ile Ser Trp Asp Ile 330 325 Gln Asp Glu Lys Asn Val Thr Cys Gln Leu Thr Phe Trp Glu Ala Ser 345 Glu Arg Thr Ile Arg Ser Glu Ala Glu Asp Ser Tyr His Phe Ser Ser 4 365 360 Ala Lys Met Thr Ala Thr Phe Leu Ser Lys Lys Gln Glu Val Asn Met 375 380 Ser Asp Ser Ala Leu Asp Cys Val Arg Asp Glu Ala Ile Asn Lys Leu 390 395 Gln Gln Ile Phe Asn Thr Ser Tyr Asn Gln Thr Tyr Glu Lys Tyr Gly 405 410 Asn Val Ser Val Phe Glu Thr Ser Gly Gly Leu Val Val Phe Trp Gln 425 Gly Ile Lys Gln Lys Ser Leu Val Glu Leu Glu Arg Leu Ala Asn Arg 435 440 Ser Ser Leu Asn Ile Thr His Arg Thr Arg Arg Ser Thr Ser Asp Asn 455 460 Asn Thr Thr His Leu Ser Ser Met Glu Ser Val His Asn Leu Val Tyr 470 475 Ala Gln Leu Gln Phe Thr Tyr Asp Thr Leu Arg Gly Tyr Ile Asn Arg 485 490 Ala Leu Ala Gln Ile Ala Glu Ala Trp Cys Val Asp Gln Arg Arg Thr 500 505 Leu Glu Val Phe Lys Glu Leu Ser Lys Ile Asn Pro Ser Ala Ile Leu 520 Ser Ala Ile Tyr Asn Lys Pro Ile Ala Ala Arg Phe Met Gly Asp Val 540 Leu Gly Leu Ala Ser Cys Val Thr Ile Asn Gln Thr Ser Val Lys Val 550 555 Leu Arg Asp Met Asn Val Lys Glu Ser Pro Gly Arg Cys Tyr Ser Arg 565 570 Pro Val Val Ile Phe Asn Phe Ala Asn Ser Ser Tyr Val Gln Tyr Gly 585 Gln Leu Gly Glu Asp Asn Glu Ile Leu Leu Gly Asn His Arg Thr Glu 600 Glu Cys Gln Leu Pro Ser Leu Lys Ile Phe Ile Ala Gly Asn Ser Ala

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615
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Tyr Glu Tyr Val Asp Tyr Leu Phe Lys Arg Met Ile Asp Leu Ser Ser
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Ile Ser Thr Val Asp Ser Met Ile Ala Leu Asp Ile Asp Pro Leu Glu
                                    650
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Asn Thr Asp Phe Arg Val Leu Glu Leu Tyr Ser Gln Lys Glu Leu Arg
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Ser Ser Asn Val Phe Asp Leu Glu Glu Ile Met Arg Glu Phe Asn Ser
                            680
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Tyr Lys Gln Arg Val Lys Tyr Val Glu Asp Lys Val Val Asp Pro Leu
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Pro Pro Tyr Leu Lys Gly Leu Asp Asp Leu Met Ser Gly Leu Gly Ala
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Ala Gly Lys Ala Val Gly Val Ala Ile Gly Ala Val Gly Gly Ala Val
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Ala Ser Val Val Glu Gly Val Ala Thr Phe Leu Lys Asn Pro Phe Gly
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Ala Phe Thr Ile Ile Leu Val Ala Ile Ala Val Val Ile Ile Thr Tyr
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Leu Ile Tyr Thr Arg Gln Arg Arg Leu Cys Thr Gln Pro Leu Gln Asn
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Leu Phe Pro Tyr Leu Val Ser Ala Asp Gly Thr Thr Val Thr Ser Gly
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Ser Thr Lys Asp Thr Ser Leu Gln Ala Pro Pro Ser Tyr Glu Glu Ser
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Val Tyr Asn Ser Gly Arg Lys Gly Pro Gly Pro Pro Ser Ser Asp Ala
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                                                    830
Ser Thr Ala Ala Pro Pro Tyr Thr Asn Glu Gln Ala Tyr Gln Met Leu
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Leu Ala Leu Ala Arg Leu Asp Ala Glu Gln Arg Ala Gln Gln Asn Gly
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Thr Asp Ser Leu Asp Gly Gln Thr Gly Thr Gln Asp Lys Gly Gln Lys
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tot ato gtt tto atg gtg otg act ttt gto aac gto ago gtg cat ota
                                                                       96
Ser Ile Val Phe Met Val Leu Thr Phe Val Asn Val Ser Val His Leu
gtg ctg agc aat ttt ccg cac ctg ggc tac ccc tgc gtc tac tat cac
                                                                      144
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Val	Leu	Ser 35	Asn	Phe	Pro	His	Leu 40	Gly	Tyr	Pro	Cys	Val 45	Tyr	Tyr	His		
							aac Asn									192	
							tta Leu									240	
							ttt Phe									288	
-	-			-		-	atg Met	-	_	-			_	_		336	
							tcg Ser 120									384	
							gac Asp									432	
							atg Met									480	
							aac Asn									528	
							ttc Phe									576	
							acg Thr 200									624	
							gtg Val									672	
							cgt Arg									720	
							tcc Ser									768	
							aag Lys									816	

260		265	270	
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-	_	Arg Thr Gly Ile S	egc tgg tcg ttc ggc Ger Trp Ser Phe Gly 100	912
			gt cgc gcc gtc cgc Cys Arg Ala Val Arg 320	960
Tyr Phe Arg Gly			ac cag gcg ctg gcc 1 'yr Gln Ala Leu Ala . 335	1008
			ac cac gac agc ttg 1 lis His Asp Ser Leu 350	
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Ser Tyr Lys Arg Ser Leu Phe Phe Phe Ser Arg Leu His Pro Lys Leu
            180
                                185
Lys Gly Thr Val Gln Phe Arg Thr Leu Ile Val Asn Leu Val Glu Val
                            200
Ala Leu Gly Phe Asn Thr Thr Val Val Ala Met Ala Leu Cys Tyr Gly
                        215
Phe Gly Asn Asn Phe Phe Val Arg Thr Gly His Met Val Leu Ala Val
                    230
                                        235
Phe Val Val Tyr Ala Ile Ile Ser Ile Ile Tyr Phe Leu Leu Ile Glu
                245
                                    250
Ala Val Phe Phe Gln Tyr Val Lys Val Gln Phe Gly Tyr His Leu Gly
                                265
Ala Phe Phe Gly Leu Cys Gly Leu Ile Tyr Pro Ile Val Gln Tyr Asp
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Thr Phe Leu Ser Asn Glu Tyr Arg Thr Gly Ile Ser Trp Ser Phe Gly
                        295
Met Leu Phe Phe Ile Trp Ala Met Phe Thr Thr Cys Arg Ala Val Arg
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                                        315
Tyr Phe Arg Gly Arg Gly Ser Gly Ser Val Lys Tyr Gln Ala Leu Ala
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                325
Thr Ala Ser Gly Glu Glu Val Ala Val Leu Ser His His Asp Ser Leu
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                                                                       96
Ala Glu Ser Ser Gly Asn Asn Ser Ser Thr Ser Thr Ser Ala Thr Thr
                                                                      144
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Ser Lys Ser Ser Ala Ser Val Ser Thr Thr Lys Leu Thr Thr Val Ala
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aca act tet gea aca act acg acg act acg acc tta teg aca act age
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Thr Thr Ser Ala Thr Thr Thr Thr Thr Thr Leu Ser Thr Thr Ser
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Thr Lys Leu Ser Ser Thr Thr His Asp Pro Asn Val Met Arg Arg His
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aac ttt act of Asn Phe Thr A	-		_		417
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Thr Lys Leu S	Ser Ser Thr 70	Thr His As	sp Pro Asn 75	Val Met Arg	Arg His 80
Ala Asn Asp A	Asp Phe Tyr 85	Lys Ala Hi	s Cys Thr	Ser His Met	Tyr Glu 95
Leu Ser Leu S	Ser Ser Phe 100	Ala Ala Tr 10		Met Leu Asn 110	Ala Leu
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		_		_	cgt Arg	_				_	-				_	192
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					ctg Leu										ctg Leu [.]	336
	-			_	aga Arg			_	_					_	_	384
-			-	_	gcc Ala	_		_			_	_	_	_	_	432
		_	_	_	ggt Gly 150	_	-						-			480
					cac His											528
					cac His											576
					ctc Leu											624
					cac His											672
					ctg Leu 230											720

					ccc Pro											768
-				-	ccc Pro			_	_							816
	-				ctc Leu	_	-		_	_		-				864
					aaa Lys											912
			-	-	ctc Leu 310	-	-	-		_		-			-	960
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					cag Gln											1344
					acg Thr											1392
cgc	caa	gaa	ctc	tac	ctc	atg	ggc	agc	ctc	gtc	cac	tcc	atg	ttg	gta	1440

Arg 465	Gln	Glu	Leu	Tyr	Leu 470	Met	Gly	Ser	Leu	Val 475	His	Ser	Met	Leu	Val 480	
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	_	gcc Ala						_	-	-		-		-		1536
		tac Tyr 515														1584
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		ttc Phe														1776
		cag Gln 595		_								-				1824
		ggc Gly														1872
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		att Ile														1968
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Thr Glu Lys His Glu Leu Leu Val Leu Val Lys Lys Ala Gln Leu Asn

285

280

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Phe 305		Tyr	Leu	Asp	Leu 310	Ser	Ala	Leu	Leu	Arg 315	Asn	Ser	Phe	His	Arg 320
	Ala	Val	Asp	Val 325	Leu	Lys	Ser	Gly	Arg 330	Cys	Gln	Met	Leu	Asp 335	Arg
Arg	Thr	Val ·	Glu 340	Met	Ala	Phe	Ala	Tyr 345	Ala	Leu	Ala	Leu	Phe 350	Ala	Ala
Ala	_	Gln 355	Glu	Glu	Ala	Gly	Thr 360	Glu	Ile	Ser	Ile	Pro 365	Arg	Ala	Leu
Asp			Ala	Ala	Leu	Leu 375	Gln	Ile	Gln	Glu	Phe 380	Met	Ile	Thr	Cys
Leu 385	Ser	Gln	Thr	Pro	Pro 390	Arg	Thr	Thr	Leu	Leu 395	Leu	Tyr	Pro	Thr	Ala 400
Val	Asp	Leu	Ala	Lys 405	Arg	Ala	Leu	Trp	Thr 410	Pro	Asp	Gln	Ile	Thr 415	Asp
Ile	Thr	Ser	Leu 420	Val	Arg	Leu	Val	Tyr 425	Ile	Leu	Ser	Lys	Gln 430	Asn	Gln
Gln	His	Leu 435	Ile	Pro	Gln	Trp	Ala 440	Leu	Arg	Gln	Ile	Ala 445	Asp	Phe	Ala
Leu	Gln 450	Leu	His	Lys	Thr	His 455	Leu	Ala	Ser	Phe	Leu 460	Ser	Ala	Phe	Ala
465			Leu	-	470		_			475					480
His	Thr	Thr	Glu	Arg 485	Arg	Glu	Ile	Phe	Ile 490	Val	Glu	Thr	Gly	Leu 495	Cys
			Glu 500					505					510		
		515	Leu				520					525			
-	530		Ser			535					540				
545			Thr		550					555					560
			Leu	565				_	570		_			575	
			Ser 580					585					590		
		595	Tyr			-	600			-		605			
	610	_	Gln			615					620				
625			Thr		630					635					640
			Ser	645			_		650	-				655	
	_	_	Asp 660					665					670		
	_	675	Val				680					685			
	690		Arg			695					700				
705			Thr	_	710					715					720
			Val	725			Ser	Ala	Ile 730	Ile	Gly	Ile	Tyr	Leu 735	Leu
Tyr	Arg	Met	Leu	Lys	Thr	Cys									

<211 <212)> 9 L> 92 2> Di 3> Hi	Ν	cyto	omega	alovi	irus											
	L> CI		. (918	3)													
cgt			gat Asp														48 .
			tca Ser 20														96
			ggc Gly													,	144
_	_		ctg Leu	_			_			-	-	-	_	-	-		192
		_	gcc Ala			_				_		_		_	-	;	240
			ttg Leu													:	288
			ttg Leu 100													:	336
			tac Tyr													:	384
_	_	_	gct Ala		_	-		_	_	-	-				_	,	432
			cgg Arg														480
			acg Thr														528
ccg	gcc	gtg	tac	acg	tgc	gtg	gac	gac	ctg	tgc	cgc	ggc	tac	gac	ctc		576

Pro Ala Val	Tyr Thr 180	Cys Val	Asp	Asp 185	Leu	Cys	Arg	Gly	Tyr 190	Asp	Leu	
acg cga ctg Thr Arg Leu 195			-			_	-					624
ttc gag ctg Phe Glu Leu 210												672
aac gaa gcc Asn Glu Ala 225												720
gct gcc gcg Ala Ala Ala												768
gtg aag gaa Val Lys Glu	_	-		_	-	-	_	_			_	816
cac cta gat His Leu Asp 275												864
cgc gtc aac Arg Val Asn 290		-	_	_					-		-	912
gct cgc taa Ala Arg 305												921
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Arg Phe Arg	5	_			10					15	_	
Ser Leu Pro	Ser His 20	Ser His	Arg	Ala 25	Ala	Asp	Leu	Met	Cys 30	Arg	Arg	
Pro Asp Cys	Gly Phe	Ser Phe	Ser 40	Pro	Gly	Pro	Val	Val 45	Leu	Leu	Trp	
Cys Cys Leu 50	Leu Leu	Pro Ile 55		Ser	Ser	Val	Ala 60	_	Ser	Val	Ala	
Pro Thr Ala	Ala Glu		Pro	Ala	Glu	Cys 75		Glụ	Leu	Thr	Arg 80	
Arg Cys Leu	_		Phe	Gln	_		Lys	Tyr	Glu			
Leu Arg Pro		Asn Val	Thr	Arg 105	90 Arg	Asp	Gly	Pro	Leu 110	95 Ser	Gln	
Leu Ile Arg	100 Tyr Arg	Pro Val	Thr		Glu	Ala	Ala	Asn		Val	Leu	

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115
                            120
Leu Asp Asp Ala Phe Leu Asp Thr Leu Ala Leu Leu Tyr Asn Asn Pro
                                            140
                        135
Asp Gln Leu Arg Ala Leu Leu Thr Leu Leu Ser Ser Asp Thr Ala Pro
                    150
                                        155
Arg Trp Met Thr Val Met Arg Gly Tyr Ser Glu Cys Gly Asp Gly Ser
                                    170
                165
Pro Ala Val Tyr Thr Cys Val Asp Asp Leu Cys Arg Gly Tyr Asp Leu
            180
                                185
Thr Arg Leu Ser Tyr Gly Arg Ser Ile Phe Thr Glu His Val Leu Gly
                            200
                                                205
Phe Glu Leu Val Pro Pro Ser Leu Phe Asn Val Val Ala Ile Arg
                        215
                                            220
Asn Glu Ala Thr Arg Thr Asn Arg Ala Val Arg Leu Pro Val Ser Thr
                    230
                                        235
Ala Ala Pro Glu Gly Ile Thr Leu Phe Tyr Gly Leu Tyr Asn Ala
                                    250
Val Lys Glu Phe Cys Leu Arg His Gln Leu Asp Pro Pro Leu Leu Arg
His Leu Asp Lys Tyr Tyr Ala Gly Leu Pro Pro Glu Leu Lys Gln Thr
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                            280
Arg Val Asn Leu Pro Ala His Ser Arg Tyr Gly Pro Gln Ala Val Asp
Ala Arg
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Met Gly Arg Lys Glu Met Met Val Arg Asp Val Pro Lys Met Val Phe
cta ata tct ata tct ttc ttg ctt gtt tct ttc ata aac tgt aaa gtt
                                                                       96
Leu Ile Ser Ile Ser Phe Leu Leu Val Ser Phe Ile Asn Cys Lys Val
             20
atq tca aaa gcg ctt tat aat cgt cct tgg agg ggc ttg gta ctg tct
                                                                      144
Met Ser Lys Ala Leu Tyr Asn Arg Pro Trp Arg Gly Leu Val Leu Ser
         35
aag ata ggc aaa tat aaa tta gat cag ctt aag tta gaa att ttg aga
                                                                      192
Lys Ile Gly Lys Tyr Lys Leu Asp Gln Leu Lys Leu Glu Ile Leu Arg
                         55
caa cta gaa acg act att tct aca aaa tac aat gta agt aaa caa ccg
                                                                      240
Gln Leu Glu Thr Thr Ile Ser Thr Lys Tyr Asn Val Ser Lys Gln Pro
                     70
                                         75
gtt aaa aat ctc act atg aac atg aca gag ttt cca caa tac tac att
                                                                      288
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Val	Lys	Asn	Leu	Thr 85	Met	Asń	Met	Thr	Glu 90	Phe	Pro	Gln	Tyr	Tyr 95	Ile	
					cag Gln			-				_			-	336
		-			ctt Leu							-			_	384
			_	_	aaa Lys	_				_				_		432
					act Thr 150											480
		_			gaa Glu			-					_			528
					gta Val							_				576
_		_	-		gta Val	-	_		_							624
		_			cgt Arg		_		_		-			_		672
					atg Met 230						_	-		_		720
				_	tta Leu				_	_				_		768
			_	_	gaa Glu			-			_	_			_	816
					tac Tyr											864
					act Thr											912
					gct Ala											960

305	31	0	31	5	320
	_		-	g tgg gtg tat r Trp Val Tyr	
-				a agc cgc aac o Ser Arg Asn 350	•
	-	-		a cta atc cgt l Leu Ile Arg 365	_
			-	g agc tcc tta t Ser Ser Leu 380	
-		l Glu Asn	_	t tcc gat agt a Ser Asp Ser 5	
•	-	_		g aga aca ttt n Arg Thr Phe	•
	-	-		t cta gat gag e Leu Asp Glu 430	_
-		_	-	c ctt acc ccg n Leu Thr Pro 445	J J
	-	-		t agc ctt tgg n Ser Leu Trp 460	
ttg cag taa Leu Gln 465					1401
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Met Ser Lys	20	r Asn Arg	25	30 g Gly Leu Val	-
35 Lys Ile Gly 50	Lys Tyr Ly	40 s Leu Asp 55	Gln Leu Ly	45 rs Leu Glu Ile 60	Leu Arg

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Gln Leu Glu Thr Thr Ile Ser Thr Lys Tyr Asn Val Ser Lys Gln Pro
                    70
Val Lys Asn Leu Thr Met Asn Met Thr Glu Phe Pro Gln Tyr Tyr Ile
                                    90
Leu Ala Gly Pro Ile Gln Asn Tyr Ser Ile Thr Tyr Leu Trp Phe Asp
                               105
           100
Phe Tyr Ser Thr Gln Leu Arg Lys Pro Ala Lys Tyr Val Tyr Ser Gln
                           120
Tyr Asn His Thr Ala Lys Thr Ile Thr Phe Arg Pro Pro Pro Cys Gly
                                           140
                       135
Thr Val Pro Ser Met Thr Cys Leu Ser Glu Met Leu Asn Val Ser Lys
                                       155
                   150
Arg Asn Asp Thr Gly Glu Gln Gly Cys Gly Asn Phe Thr Thr Phe Asn
               165
                                   170
Pro Met Phe Phe Asn Val Pro Arg Trp Asn Thr Lys Leu. Tyr Val Gly
                               185
Pro Thr Lys Val Asn Val Asp Ser Gln Thr Ile Tyr Phe Leu Gly Leu
                            200
Thr Ala Leu Leu Arg Tyr Ala Gln Arg Asn Cys Thr His Ser Phe
                        215
                                           220
Tyr Leu Val Asn Ala Met Ser Arg Asn Leu Phe Arg Val Pro Lys Tyr
                   230
                                       235
Ile Asn Gly Thr Lys Leu Lys Asn Thr Met Arg Lys Leu Lys Arg Lys
               245
                                    250
Gln Ala Pro Val Lys Glu Gln Phe Glu Lys Lys Ala Lys Lys Thr Gln
           260
                               265
Ser Thr Thr Thr Pro Tyr Phe Ser Tyr Thr Thr Ser Ala Ala Leu Asn
                           280
Val Thr Thr Asn Val Thr Tyr Ser Ile Thr Thr Ala Ala Arg Arg Val
                        295
Ser Thr Ser Thr Ile Ala Tyr Arg Pro Asp Ser Ser Phe Met Lys Ser
                    310
                                        315
Ile Met Ala Thr Gln Leu Arg Asp Leu Ala Thr Trp Val Tyr Thr Thr
                325
                                    330
Leu Arg Tyr Arg Gln Asn Pro Phe Cys Glu Pro Ser Arg Asn Arg Thr
                               345
Ala Val Ser Glu Phe Met Lys Asn Thr His Val Leu Ile Arg Asn Glu
                           360
                                                365
Thr Pro Tyr Thr Ile Tyr Gly Thr Leu Asp Met Ser Ser Leu Tyr Tyr
                        375
                                           380
Asn Glu Thr Met Phe Val Glu Asn Lys Thr Ala Ser Asp Ser Asn Lys
                   390
                                       395
Thr Thr Pro Thr Ser Pro Ser Met Gly Phe Gln Arg Thr Phe Ile Asp
               405
                                   410
Pro Leu Trp Asp Tyr Leu Asp Ser Leu Leu Phe Leu Asp Glu Ile Arg
                                425
Asn Phe Ser Leu Arg Ser Pro Thr Tyr Val Asn Leu Thr Pro Pro Glu
                           440
                                               445
His Arg Arg Ala Val Asn Leu Ser Thr Leu Asn Ser Leu Trp Trp
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Leu Gln
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		Ile Leu Val	tcg cag tac acc Ser Gln Tyr Thi	
Ser Thr Pro Cy 65			ctg cag gtg cag Leu Gln Val Glr 75	
			tcg gtc aac gtc Ser Val Asn Val	
	g Ser Ile Cys		gag ccc atg tcc Glu Pro Met Ser 110	: Ile Tyr
			atc ccc agc atc Ile Pro Ser Ile 125	
His His Tyr Pr		Glu Arg Lys	cac cga cac cto His Arg His Lev 140	
			cag atg tgg cag Gln Met Trp Glr 155	
			cag cag aac cag Gln Gln Asn Glr	
	Tyr Tyr Thr		gtg ttt ccc acc Val Phe Pro Thi 190	Lys Asp
			gag ctg gtt tgg Glu Leu Val Cys 205	

		cgc Arg										672
		ctg Leu										720
		gtc Val										768
	-	ccg Pro 260			_	_			-			816
		tgt Cys										864
		ctg Leu										912
_	_	aag Lys	_	_		_	_			_	_	960
		cag Gln										1008
		cgt Arg 340										1056
		ctg Leu										1104
		cag Gln										1152
		cac His										1200
		tcg Ser										1248
		gtc Val 420										1296

gcg ggc cgc Ala Gly Arg 435							1344
ggc gtt atg Gly Val Met 450					_	-	1392
gaa gag gac Glu Glu Asp 465		ı Asp Ser					1440
gtg ttc acc Val Phe Thr							1488
gtg ccc atg Val Pro Met							1536
ttc ttc tgg Phe Phe Trp 515		_	_				1584
ggc gta tgg Gly Val Trp 530							1632
gac gcc ttg Asp Ala Leu 545		Cys Ile					1680
ggt tga Gly							1686
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Pro Val Leu	20	-	25		30	-	
35 Arg Val Ser		40			45		
50 Ser Thr Pro		55		60	_	_	
65 Tyr Phe Thr	70			75		80	
Pro Thr Gly	85 Arg Ser Il 100	e Cys Pro	90 Ser Gln 105	Glu Pro	Met Ser 110	95 Ile Tyr	

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Val Tyr Ala Leu Pro Leu Lys Met Leu Asn Ile Pro Ser Ile Asn Val
                            120
                                                125
His His Tyr Pro Ser Ala Ala Glu Arg Lys His Arg His Leu Pro Val
                        135
                                            140
Ala Asp Ala Val Ile His Ala Ser Gly Lys Gln Met Trp Gln Ala Arg
                    150
                                        155
Leu Thr Val Ser Gly Leu Ala Trp Thr Arg Gln Gln Asn Gln Trp Lys
                                   170
Glu Pro Asp Val Tyr Tyr Thr Ser Ala Phe Val Phe Pro Thr Lys Asp
           180
                                185
Val Ala Leu Arg His Val Val Cys Ala His Glu Leu Val Cys Ser Met
                            200
Glu Asn Thr Arg Ala Thr Lys Met Gln Val Ile Gly Asp Gln Tyr Val
                       215
                                            220
Lys Val Tyr Leu Glu Ser 'Phe Cys Glu Asp Val Pro Ser Gly Lys Leu
                    230
                                       235
Phe Met His Val Thr Leu Gly Ser Asp Val Glu Glu Asp Leu Thr Met
                                    250
                245
Thr Arg Asn Pro Gln Pro Phe Met Arg Pro His Glu Arg Asn Gly Phe
                                265
            260
Thr Val Leu Cys Pro Lys Asn Met Ile Ile Lys Pro Gly Lys Ile Ser
                           280
His Ile Met Leu Asp Val Ala Phe Thr Ser His Glu His Phe Gly Leu
                        295
                                            300
Leu Cys Pro Lys Ser Ile Pro Gly Leu Ser Ile Ser Gly Asn Leu Leu
                    310
                                        315
Met Asn Gly Gln Gln Ile Phe Leu Glu Val Gln Ala Ile Arg Glu Thr
               325
                                    330
Val Glu Leu Arg Gln Tyr Asp Pro Val Ala Ala Leu Phe Phe Phe Asp
                                345
Ile Asp Leu Leu Gln Arg Gly Pro Gln Tyr Ser Glu His Pro Thr
                            360
Phe Thr Ser Gln Tyr Arg Ile Gln Gly Lys Leu Glu Tyr Arg His Thr
                        375
Trp Asp Arg His Asp Glu Gly Ala Ala Gln Gly Asp Asp Asp Val Trp
                   390
                                       395
Thr Ser Gly Ser Asp Ser Asp Glu Glu Leu Val Thr Thr Glu Arg Lys
                405
                                    410
Thr Pro Arg Val Thr Gly Gly Gly Ala Met Ala Gly Ala Ser Thr Ser
            420
                                425
Ala Gly Arg Lys Arg Lys Ser Ala Ser Ser Ala Thr Ala Cys Thr Ser
       435
                            440
                                                445
Gly Val Met Thr Arg Gly Arg Leu Lys Ala Glu Ser Thr Val Ala Pro
                        455
                                            460
Glu Glu Asp Thr Asp Glu Asp Ser Asp Asn Glu Ile His Asn Pro Ala
                    470
                                        475
Val Phe Thr Trp Pro Pro Trp Gln Ala Gly Ile Leu Ala Arg Asn Leu
                                    490
Val Pro Met Val Ala Thr Val Gln Gly Gln Asn Leu Lys Tyr Gln Glu
            500
                                505
                                                    510
Phe Phe Trp Asp Ala Asn Asp Ile Tyr Arg Ile Phe Ala Glu Leu Glu
                            520
Gly Val Trp Gln Pro Ala Ala Gln Pro Lys Arg Arg His Arg Gln
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Asp Ala Leu Pro Gly Pro Cys Ile Ala Ser Thr Pro Lys Lys His Arg
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Gly
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gtc aac ttt ctg cgc cat ctc acg caa aag ccc gac gtg gat ctc gag
                                                                        96
Val Asn Phe Leu Arg His Leu Thr Gln Lys Pro Asp Val Asp Leu Glu
             20
                                                      30
                                                                      144
gca cac ccc aag atc ctg aaa aaa tgt ggc gaa aaa cgc ctg cac cgg
Ala His Pro Lys Ile Leu Lys Lys Cys Gly Glu Lys Arg Leu His Arg
         35
cgt acg gtg ctg ttc aac gag ctc atg ctt tgg ttg gga tac tac cgc
                                                                      192
Arg Thr Val Leu Phe Asn Glu Leu Met Leu Trp Leu Gly Tyr Tyr Arg
gag ctg cgt ttt cac aac ccc gac ctc tcc tca gtg ctc gag gag ttc
                                                                      240
Glu Leu Arg Phe His Asn Pro Asp Leu Ser Ser Val Leu Glu Glu Phe
gag gtg cgt tgc gtg gcc gtg gcg cgt cgc ggc tac act tac ccg ttc
                                                                      288
Glu Val Arg Cys Val Ala Val Ala Arg Arg Gly Tyr Thr Tyr Pro Phe
ggt gat cgt ggt aag gcg cgt gac cac ctg gct gtg cta gac cgt acc
                                                                      336
Gly Asp Arg Gly Lys Ala Arg Asp His Leu Ala Val Leu Asp Arg Thr
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                                 105
                                                     110
gaa ttc gat acg gac gtg cgc cac gat gcc gag atc gtg gaa cgc gcg
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Glu Phe Asp Thr Asp Val Arg His Asp Ala Glu Ile Val Glu Arg Ala
        115
                            120
ctc gta agc gcg gtc att ctg gcc aag atg tcg gtg cgc gag acg ctg
                                                                      432
Leu Val Ser Ala Val Ile Leu Ala Lys Met Ser Val Arg Glu Thr Leu
    130
                        135
gtc aca gcc atc ggc cag acg gaa ccc atc gcc ttt gtg cac ctc aag
                                                                       480
Val Thr Ala Ile Gly Gln Thr Glu Pro Ile Ala Phe Val His Leu Lys
145
                    150
                                         155
                                                             160
gat acg gag gtg cag cgc att gaa gaa aac ctg gag ggt gtg cgc cgt
                                                                      528
Asp Thr Glu Val Gln Arg Ile Glu Glu Asn Leu Glu Gly Val Arg Arg
                165
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aac atg ttc tgc gtg aaa ccg ctc gac ctt aac ctg gac cgg cac gcc
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Asn	Met	Phe	Cys 180	Val	Lys	Pro	Leu	Asp 185	Leu	Asn	Leu	Asp	Arg 190	His	Ala	
	-		_	_	aac Asn	_	_		_				_		-	624
					cgc Arg											672
_	, ,			_	gag Glu 230	_	_	_	_	_	_	_		_	_	720
_	_				gat Asp				_	_				_		768
-		_			gac Asp	_	_	-	_	-	-		-			816
	-		-		tac Tyr		-	-			~	-			_	864
					tcg Ser											912
		_			ggc Gly 310		_	_		-	_	_		_		960
	-		_	_	ccg Pro	_	_			_			_	_		1008
					ccg Pro											1056
					cac His											1104
					cgt Arg											1152
					aaa Lys 390											1200
					cgt Arg											1248

				405					410					415			
					agg Arg												1296
_	-	-	_	_	gag Glu					_			_		_	:	1344
					ggc Gly											:	1392
					tcc Ser 470												1440
			_		caa Gln		_	-			-		-		-		1488
	-	_			aag Lys	-	_	_			_	_			_	:	1536
					ccg Pro												1584
	-		-		gac Asp	-	_	_						_			1632
					gac Asp 550												1680
					gtc Val												1728
					gga Gly												1776
					cct Pro												1824
					gcg Ala												1872
					gcg Ala 630												1920

	ccg Pro															1968
	ccc Pro								_	-						2016
	acc Thr					_	_			_	_					2064
	gcg Ala 690													-		2112
	gca Ala															2160
_	gac Asp				-	-		_					_	_		2208
_	tac Tyr			_	-		_			_	_	_			_	2256
-	acg Thr	_		-		-		-	-		_			_	-	2304
	aca Thr 770															2352
_	cac His			-	_	-	_	-		_	_	_	_	_	_	2400
	agg Arg				-	-			_	-	_	_	_		-	2448
_	ctg Leu	_				_	_	_				_				2496
	gtg Val															2544
_	tca Ser 850	-	-			_	-	-		_		_	_	_		2592

	ccc Pro	-		_	_	_		-		_		_		_		2640
	gcc Ala															2688
	ggt Gly															2736
	cga Arg	_		-	_			-		-		-		_	_	2784
	ggc Gly 930															2832
	acc Thr	-			_	_		_	_		_	_	_	_		2880
	ccg Pro															2928
-	ttg Leu			_	_	_	_			-	-	_				2976
	aag Lys		Thr					Thr					Leu			3024
	cag Gln 1010	Lys	_		-		Pro		-			Āla		_		3072
	gcc Ala 5					Asp					Ile					3120
	aag Lys		_		Thr		_	tag								3147

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<211> 1048

<212> PRT

<213> Human cytomegalovirus

<400> 16

Met Ser Leu Gln Phe Ile Gly Leu Gln Arg Arg Asp Val Val Ala Leu $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Val Asn Phe Leu Arg His Leu Thr Gln Lys Pro Asp Val Asp Leu Glu Ala His Pro Lys Ile Leu Lys Lys Cys Gly Glu Lys Arg Leu His Arg Arg Thr Val Leu Phe Asn Glu Leu Met Leu Trp Leu Gly Tyr Tyr Arg 55 Glu Leu Arg Phe His Asn Pro Asp Leu Ser Ser Val Leu Glu Glu Phe Glu Val Arg Cys Val Ala Val Ala Arg Arg Gly Tyr Thr Tyr Pro Phe 90 Gly Asp Arg Gly Lys Ala Arg Asp His Leu Ala Val Leu Asp Arg Thr 100 105 Glu Phe Asp Thr Asp Val Arg His Asp Ala Glu Ile Val Glu Arg Ala 120 125 Leu Val Ser Ala Val Ile Leu Ala Lys Met Ser Val Arg Glu Thr Leu 135 140 Val Thr Ala Ile Gly Gln Thr Glu Pro Ile Ala Phe Val His Leu Lys 150 155 Asp Thr Glu Val Gln Arg Ile Glu Glu Asn Leu Glu Gly Val Arg Arg 165 170 Asn Met Phe Cys Val Lys Pro Leu Asp Leu Asn Leu Asp Arg His Ala 185 Asn Thr Ala Leu Val Asn Ala Val Asn Lys Leu Val Tyr Thr Gly Arg 200 205 Leu Ile Met Asn Val Arg Arg Ser Trp Glu Glu Leu Glu Arg Lys Cys 215 220 Leu Ala Arg Ile Gln Glu Arg Cys Lys Leu Leu Val Lys Glu Leu Arg 230 235 Met Cys Leu Ser Phe Asp Ser Asn Tyr Cys Arg Asn Ile Leu Lys His 250 Ala Val Glu Asn Gly Asp Ser Ala Asp Thr Leu Leu Glu Leu Leu Ile 265 Glu Asp Phe Asp Ile Tyr Val Asp Ser Phe Pro Gln Ser Ala His Thr 280 Phe Leu Gly Ala Arg Ser Pro Ser Leu Glu Phe Asp Asp Asp Ala Asn 295 300 Leu Leu Ser Leu Gly Gly Gly Ser Ala Phe Ser Ser Val Pro Lys Lys 310 315 His Val Pro Thr Gln Pro Leu Asp Gly Trp Ser Trp Ile Ala Ser Pro 325 330 Trp Lys Gly His Lys Pro Phe Arg Phe Glu Ala His Gly Ser Leu Ala 340 345 Pro Ala Ala Glu Ala His Ala Ala Arg Ser Ala Ala Val Gly Tyr Tyr 360 Asp Glu Glu Glu Lys Arg Arg Glu Arg Gln Lys Arg Val Asp Asp Glu 375 380 Val Val Gln Arg Glu Lys Gln Gln Leu Lys Ala Trp Glu Glu Arg Gln 390 395 Gln Asn Leu Gln Gln Arg Gln Gln Gln Pro Pro Pro Ala Arg Lys 405 410 Pro Ser Ala Ser Arg Arg Leu Phe Gly Ser Ser Ala Asp Glu Asp Asp 425 Asp Asp Asp Asp Glu Lys Asn Ile Phe Thr Pro Ile Lys Lys Pro 440 445 Gly Thr Ser Gly Lys Gly Ala Ala Ser Gly Gly Gly Val Ser Ser Ile 455 460 Phe Ser Gly Leu Leu Ser Ser Gly Ser Gln Lys Pro Thr Ser Gly Pro

465					470					475					480
Leu	Asn	Ile	Pro	Gln 485	Gln	Gln	Gln	Arg	His 490	Ala	Ala	Phe	Ser	Leu 495	Val
Ser	Pro	Gln	Val 500	Thr	Lys	Ala	Ser	Pro 505	Gly	Arg	Val	Arg	Arg 510	Asp	Ser
Ala	Trp	Asp 515	Val	Arg	Pro	Leu	Thr 520	Glu	Thr	Arg	Gly	Asp 525	Leu	Phe	Ser
Gly	Asp 530	Glu	Asp	Ser	Asp	Ser 535	Ser	Asp	Gly	Tyr	Pro 540	Pro	Asn	Arg	Gln
545			Phe		550					555		_			560
		-	Pro	565					570	_				575	
			Gly 580		_			585			_		590		
		595	Pro				600					605		•	
	610		Thr		•	615					620		_		
625	_		Pro		630				_	635					640
			Glu	645					650					655	
			Arg 660					665					670		
		675·		_	_		680			-		685			
	690		Arg			695					700				
705			Ser	•	710					715		_	_	_	720
	_		Gly	725					730					735	
_	_		Asn 740	_				745					750		
		755	Ser				760					765			
	770		Thr			775				_	780	_			
785	птъ	Ата	Ser	Ата	790	Ala	GIII	1111	val	795	ser	IIIL	PIO	vai	800
_	_		Glu	805				-	810					815	
			Gln 820					825		_			830		
		835	Ser				840					845			
	850		Ser			855					860	-	-		
865			Thr		870					875					880
			Ser	885					890					895	
	_		Pro 900			_		905		-		-	910		
GLY	Arg	Pro 915	Pro	Ser	val	Pro	Val 920	Ser	GTÀ	Ser	Ala	Pro 925	GLY	Arg	Leu

Ser Gly Ser Ser Arg Ala Ala Ser Thr Thr Pro Thr Tyr Pro Ala Val 935 Thr Thr Val Tyr Pro Pro Ser Ser Thr Ala Lys Ser Ser Val Ser Asn 950 955 Ala Pro Pro Val Ala Ser Pro Ser Ile Leu Lys Pro Gly Ala Ser Ala 965 970 Ala Leu Gln Ser Arg Ser Thr Gly Thr Ala Ala Val Gly Ser Pro 985 Val Lys Ser Thr Thr Gly Met Lys Thr Val Ala Phe Asp Leu Ser Ser 1005 1000 Pro Gln Lys Ser Gly Thr Gly Pro Gln Pro Gly Ser Ala Gly Met Gly 1015 1020 Gly Ala Lys Thr Pro Ser Asp Ala Val Gln Asn Ile Leu Gln Lys Ile 1030 · 1035 Glu Lys Ile Lys Asn Thr Glu Glu 1045